



New Link, 0003.ST25
SEQUENCE LISTING

<110> Link, Charles
<120> Methods and Compositions for Elucidating Protein Expression
Profiles in Cells
<130> 05237.0003.CPUS00
<140> US 10/660,893
<141> 2003-09-12
<160> 5
<170> PatentIn version 3.2
<210> 1
<211> 9
<212> PRT
<213> Artificial
<220>
<223> Synthesized in laboratory
<400> 1

Tyr Pro Tyr Asp Val Pro Asp Tyr Ala
1 5

<210> 2
<211> 10
<212> PRT
<213> Artificial
<220>
<223> Synthesized in laboratory
<400> 2

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
1 5 10

<210> 3
<211> 8
<212> PRT
<213> Artificial
<220>
<223> Synthesized in laboratory
<400> 3

Asp Tyr Lys Asp Asp Asp Asp Lys
1 5

<210> 4
<211> 375
<212> DNA
<213> Homo sapiens

<220>
 <221> misc_feature
 <222> (3)..(4)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (8)..(8)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (350)..(350)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (358)..(374)
 <223> n is a, c, g, or t

<400> 4	
ttnnccgnga aagctcctcg cccttgctca ccatgggatg ccatttccta ggtctgcctc	60
ttggccgttt ttctccaatg gtctctgctt tcttctgggc tgcttttagag gggctcttgt	120
ttttgctgcc tttgggtctt cctctggggtc tcttaggaga gggctcacag gttggctctt	180
gctgctgctt cctggggtcgg ccgcgtcctc gcttctgtgg caccggggcg gcagggtgtc	240
cctgggctga tgtggacggc tgcccggcgc cctcaccgcg tgcgctcatc ctgcctcccc	300
ccgccgtac cactgcctct cttttttttt tttttttttt tttttgaaan ccccggnnn	360
nnnnnnnnnnn nnnnc	375

<210> 5
 <211> 333
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (3)..(3)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (11)..(11)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (106)..(106)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (116)..(116)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (168)..(168)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (179)..(179)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (204)..(204)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (221)..(221)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (224)..(224)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (231)..(231)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (254)..(254)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (272)..(272)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (275)..(275)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (282)..(282)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (285)..(286)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (289)..(289)
 <223> n is a, c, g, or t

<220>
 <221> misc_feature
 <222> (292)..(293)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (296)..(296)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (299)..(299)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (301)..(301)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (304)..(304)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (306)..(308)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (311)..(311)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (326)..(327)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (329)..(329)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (331)..(331)

<223> n is a, c, g, or t

<220>

<221> misc_feature

<222> (333)..(333)

<223> n is a, c, g, or t

<400> 5

tcn'gcgacca nctcctcgcc cttgctcacc atgggatgct cccggtggtg ggtcgggtggt 60

ccctgggcag gggctctcaa atcccggacg agccccaaa tgaaanaccc ccgtcntggg 120

tagtcaatca ctcaaggag accctcccaa ggaacagcga gaccactntt cggatgcana 180

cagcaagagg ctttattggg aatncgggta cccggg'gcac ncantctatc ngaagactgg 240

cgttattttt tttntttttt ttttttgaat tncnggggac anccnnctna gnntanctnc 300

nctntnnnct nccctcctta cttctnntnt ntn

333